Changed a file from non-ASCII to ASCII Changed the margins in cases where the sequence text was "wrapped" down to the aead fine and fine application are to the current Application Data section, specifically: Edited a formal error in the Current Application Data section, specifically: Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or the current Application Data. Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer Changed the spelling of a mandatory field (the headings or subheadings), specifically; Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were edited were: Inserted or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited: Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted extra, invalid, headings used by an applicant, specifically: Corrected an obvious error in the beginning/end of bles; secretary initials/filename at end or page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be defeted.	Changed a file from non-ASCII to ASCII Changed the margins in cases where the sequence text was "wrapped" down to the agest line. It is felled by: Conflect by:
Changed a file from non-ASCII to ASCII Changed the margins in cases where the sequence text was "wrapped" down to the aexidine per Edited a format error in the Current Application Data section, specifically: Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or the other application Data. Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer Changed the spelling of a mandatory field (the headings or subheadings), specifically; Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheadings, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an orror in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.	Changed a file from non-ASCII to ASCII Changed the margins in cases where the sequence text was "wrapped" down to the applicant on the Current Application Data section, specifically: Edited a format error in the Current Application Data section, specifically: Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other Added the mandatory heading and subheadings for "Current Application Data". Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer. Changed the spelling of a mandatory field (the headings or subheadings), specifically; Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were edited were: Inserted or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited: Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted Extra, invalid, headings used by an applicant, specifically: Deleted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted. Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length: field accordingly (error due to a Patentin bug). Sequences corrected:
Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other	
Edited a format error in the Current Application Data section, specifically: Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other Added the mandatory heading and subheadings for 'Current Application Data'. Edited the 'Number of Sequences' field. The applicant spelled out a number instead of using an integer Changed the spelling of a mandatory field (the headings or subheadings), specifically; Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were edited were: Inserted or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited: Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted:	Edited a format error in the Current Application Data section, specifically: Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other Added the mandatory heading and subheadings for 'Current Application Data'. Edited the 'Number of Sequences' field. The applicant spelled out a number instead of using an integer. Changed the spelling of a mandatory field (the headings or subheadings), specifically; Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were edited were: Inserted or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited: Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted: one-ASCII 'garbage' at the beginning/end of files: secretary initials/filename at end of life page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A 'Hard Page Break' code was inserted by the applicant. All occurrences had to be defeted. Deleted ending stop codon in amino acid sequences and adjusted the '(A)Length: 'field accordingly (error due to a Patentin bug). Sequences corrected:
Edited the Current Application Data section with the actual current number. The number inputted by the applicant was some the prior application data; or some other some of the prior application data; or some other some of the prior application data; or some other some of the prior application data; or some other some of the prior application data; or some other some of the prior application data; or some other some of the prior application data; or some other some of the prior application data; or some other some of the prior and the prior and the prior and the prior application data; or some other some of the prior and the prior and the prior application data; or some other some other some of the prior application data; or some other some other some other some other data and the prior application data and the prior applicati	Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other Added the mandatory heading and subheadings for "Current Application Data". Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer. Changed the spelling of a mandatory field (the headings or subheadings), specifically: Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted mandatory headings, specifically: Corrected an obvious erro: in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an orror in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted. Deleted ending step codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentin bug). Sequences corrected:
Added the mandatory heading and subheadings for "Current Application Data". Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer Changed the spelling of a mandatory field (the headings or subheadings), specifically; Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted mandatory headings, specifically: Corrected mandatory headings, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.	Added the mandatory heading and subheadings for "Current Application Data". Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer. Changed the spelling of a mandatory field (the headings or subheadings), specifically; Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were edited were: Inserted or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited: Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII garbage at the beginning/end of files: secretary initials/filename at end of file page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A 'Hard Page Break' code was inserted by the applicant. All occurrences had to be defeted. Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length: field accordingly (error due to a Patentin bug). Sequences corrected:
Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer Changed the spelling of a mandatory field (the headings or subheadings), specifically; Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were edited were: Inserted or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited: Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted Edit one-ASCII garbage at the beginning/end of files; secretary initials/filename at end of page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically:	Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer. Changed the spelling of a mandatory field (the headings or subheadings), specifically; Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted: Innon-ASCII garbage at the beginning/end of files; secretary initials/filename at end of file page numbers throughout text; the headings is specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A 'Hard Page Break' code was inserted by the applicant. All occurrences had to be deleted. Deleted ending stop codon in amino acid sequences and adjusted the '(A)Length:' field accordingly (error due to a PatentIn bug). Sequences corrected:
Changed the spelling of a mandatory field (the headings or subheadings), specifically: Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were edited were: Inserted or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited: Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII garbage at the beginning/end of files; secretary initials/filename at end of page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Ediled identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically:	Changed the spelling of a mandatory field (the headings or subheadings), specifically: Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII 'garbage' at the beginning/end of files; secretary initials/filename at end of file page numbers throughout text; other invalid text, such as secretary initials/filename at end of file corrected an obvious error in the response, specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A 'Hard Page Break' code was inserted by the applicant. All occurrences had to be deleted. Deleted ending stop codon in amino acid sequences and adjusted the '(A)Length:' field accordingly (error due to a Patentin bug). Sequences corrected:
Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII garbage at the beginning/end of files; secretary initials/filename at end of page numbers throughout text; other invalid text, such as linserted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically:	Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII garbage at the beginning/end of files; secretary initials/filename at end of file page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A 'Hard Page Break' code was inserted by the applicant. All occurrences had to be deleted. Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length: field accordingly (error due to a Patentin bug). Sequences corrected:
Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII garbage at the beginning/end of files; secretary initials/filename at end of page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Ediled identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically:	Inserted or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited: Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII 'garbage' at the beginning/end of files; secretary initials/filename at end of file page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A 'Hard Page Break' code was inserted by the applicant. All occurrences had to be deleted. Deleted endIng stop codon in amino acid sequences and adjusted the '(A)Length:' field accordingly (error due to a Patentin bug). Sequences corrected:
Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted: Inon-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically: Corrected an obvious erro: in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.	Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII garbage at the beginning/end of files; secretary initials/filename at end of file page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted. Deleted endIng stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentth bug). Sequences corrected:
applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted: Inon-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.	Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII garbage at the beginning/end of files; secretary initials/filename at end of file page numbers throughout text; other invalid text, such as secretary initials/filename at end of file corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted. Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentin bug). Sequences corrected:
Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.	Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII "garbage" at the beginning/end of files: secretary initials/filename at end of file page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically: Corrected an obvious erro: in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted. Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected:
Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.	Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of file page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted. Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected:
page numbers throughout text; other invalid text, such as	Inserted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted. Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentin bug). Sequences corrected:
Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.	Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted. Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentin bug). Sequences corrected:
Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.	Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted. Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected:
Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.	Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted. Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentin bug). Sequences corrected:
Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.	Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted. Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentin bug). Sequences corrected:
A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.	A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted. Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentin bug). Sequences corrected:
	Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentin bug). Sequences corrected:
Date to the state of the state	due to a PatentIn bug). Sequences corrected:
due to a Patentin bug). Sequences corrected:	Other
Other:	=

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

....

OIPE

RAW SEQUENCE LISTING

DATE: 12/12/2001

PATENT APPLICATION: US/10/005,200

TIME: 14:19:50

Input Set : A:\Sequences for submission.txt Output Set: N:\CRF3\12112001\I005200.raw

Does Not Comply Corrected Diskette Needed

4 <110> APPLICANT: DeltaGen, Inc.

6 <120> TITLE OF INVENTION: TRANSGENIC MICE CONTAINING RPTPB

TYROSINE PHOSPHATASE GENE DISRUPTIONS

10 <130> FILE REFERENCE: 741 PCT

C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/005,200

C--> 12 <141> CURRENT FILING DATE: 2001-12-04

12 <150> PRIOR APPLICATION NUMBER: US 60/251,897

13 <151> PRIOR FILING DATE: 2000-12-06

15 <150> PRIOR APPLICATION NUMBER: US 60/302,260

16 <151> PRIOR FILING DATE: 2001-06-28

18 <160> NUMBER OF SEQ ID NOS: 3

20 <170> SOFTWARE: FastSEQ for Windows Version 4.0

ERRORED SEQUENCES

- 73 <210> SEQ ID NO: 3
- 74 <211> LENGTH: 200
- 75 <212> TYPE: DNA
- 76 <213> ORGANISM: Artificial Sequence
- 78 <220> FEATURE:
- 79 <223> OTHER INFORMATION: Targeting Vector
- 81 <400> SEQUENCE: 3
- 82 ggatgagttg aagccagaac agcagcaccc tctcccttcc tacctggagt acagacacaa 60
- 83 egectecate egagtetace agaceaatta ttttgecage aaatgtgetg aaagteeega 120
- 84 cagcagttct aaaagtttca acattaagct tggagcagag atggacagcc tcggtggcaa 180
- 85 atqtqatccc agtcagcaga

200

E--> 86(1

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/005,200

DATE: 12/12/2001

TIME: 14:19:51

Input Set : A:\Sequences for submission.txt
Output Set: N:\CRF3\12112001\1005200.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:86 M:254 E: No. of Bases conflict, LENGTH:Input:1 Counted:200 SEQ:3

DATE: 12/17/2001

OIPE

PATENT APPLICATION: US/10/005,200 TIME: 12:48:31 Input Set : A:\PTO.DC.txt Output Set: N:\CRF3\12172001\J005200.raw 4 <110> APPLICANT: DeltaGen, Inc. 6 <120> TITLE OF INVENTION: TRANSGENIC MICE CONTAINING RPTPB TYROSINE PHOSPHATASE GENE DISRUPTIONS 10 <130> FILE REFERENCE: 741 PCT C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/005,200

C--> 12 <141> CURRENT FILING DATE: 2001-12-04 12 <150> PRIOR APPLICATION NUMBER: US 60/251,897 13 <151> PRIOR FILING DATE: 2000-12-06 15 <150> PRIOR APPLICATION NUMBER: US 60/302,260 16 <151> PRIOR FILING DATE: 2001-06-28 18 <160> NUMBER OF SEQ ID NOS: 3 20 <170> SOFTWARE: FastSEQ for Windows Version 4.0 22 <210> SEQ ID NO: 1 23 <211> LENGTH: 1752 24 <212> TYPE: DNA 25 <213> ORGANISM: Mus musculus 27 <400> SEQUENCE: 1 28 gtcaaggaag aggtacctgg tgtccatcaa ggtgcagtcg gccggcatga ccagtgaggt 60 29 ggttgaagat agcaccatca ccatgataga ccgcccgcct caaccgcctc cacacatccg 120 30 tgtgaatgaa aaggatgtgc taatcagcaa atcttccatc aactttactg tcaactgcag 180 31 ctggttcagc gacaccaacg gagcggttgg gtactttgct gtggtggtga gagaggccga 240 32 cagcatggat gagttgaagc cagaacagca gcaccctctc ccttcctacc tggagtacag 300 33 acacaacgcc tccatccgag tctaccagac caattatttt gccagcaaat gtgctgaaag 360 34 tocogacago agttotaaaa gtttoaacat taagottgga goagagatgg acagootogg 420 35 tggcaaatgt gatcccagtc agcagaaatt ctgtgatgga ccgctgttgc cacacaccgc 480 36 ctacagaatc agcatccggg cttttacaca gctatttgac gaggacttga aagagttcac 540 37 caaacctctc tactcggata cgttcttctc tatgcccatc accacagagt cagagccctt 600 38 gtttggagtt attgaaggtg tgagtgctgg cctgtttcta attggcatgc tggtggccct 660 39 tgttgccttc ttcatctgca gacagaaagc tagccacagc agggaaaggc catctgcccg 720 40 gctcagcatt cgtagggacc ggcctttgtc tgtccatctg aatctgggcc agaaaggcaa 780 41 coggaaaact tottgoocca taaagatcaa toagtttgaa gggoatttoa tgaagotgoa 840 42 ggcagactcc aactaccttc tatccaagga atatgaggac ttaaaagacg tgggtagaag 900 43 ccagtcatgt gacattgccc tcttgcctga gaatcgaggg aaaaatcgat acaacaacat 960 44 attgeettat gatgeeteaa gagtgaaget etegaatgte gatgaegaee ettgetetga 1020 45 ctacatcaac gccagctaca tccccggtaa caacttcaga cgagaataca tcgccactca 1080 46 gggaccgctt ccaggcacca aggatgactt ctggaagatg gcgtgggagc agaacgttca 1140 47 caacatcgtc atggtgaccc agtgtgttga aaagggccga gtgaagtgtg accattactg 1200 48 gccagcagac caggaccccc tctactacgg tgatctcatc ctacagatgg tctcggagtc 1260 49 cgtgctcccc gagtggacca tcagggagtt taagatatgc agtgaagaac agttggatgc 1320 50 acacagacte atecgteact tteactacae ggtgtggeea gaccatgggg teccagagae 1380 51 cacccagtee etgatecaat ttgtgaggae agteagggae tacateaaca gaageeeegg 1440 52 ggctgggccc tccgtagtgc actgcagcgc tggtgtgggc agaacaggga cgttcgttgc 1500 53 cctggaccgg atcctccagc agttggactc taaggactcc gtggacattt atggggcagt 1560 54 gcatgaccta agactccaca gggttcacat ggtccagacc gagtgtcaat atgtgtatct 1620

55 gcatcagtgt gtaagagacg tctcagagca aagaaactgc ggaaacgagc aagagaaagg 1680 56 gggtgtttcg atttatgaga atgtgaatca gagtatcaca gagatgcaat ctactcgaga 1740

RAW SEQUENCE LISTING

57 cattaagaat tc

1752



RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/005,200

DATE: 12/17/2001 TIME: 12:48:31

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\12172001\J005200.raw

- 59 <210> SEQ ID NO: 2
- 60 <211> LENGTH: 200
- 61 <212> TYPE: DNA
- 62 <213> ORGANISM: Artificial Sequence
- 64 <220> FEATURE:
- 65 <223> OTHER INFORMATION: Targeting Vector
- 67 <400> SEQUENCE: 2
- 68 geogeoceca gaactecaeg geoattgeet getettggat accteetgae teegactttg 60
- 69 atggctacag cattgagtgc cgaaaaatgg atacccaaga aatcgagttt tccagaaagc 120
- 70 tggagaaaga aaaatcactg ctcaacatca tgatgttagt acctcataag aggtacctgg 180
- 71 tgtccatcaa ggtgcagtcg
- 73 <210> SEQ ID NO: 3
- 74 <211> LENGTH: 200 75 <212> TYPE: DNA
- 76 <213> ORGANISM: Artificial Sequence
- 78 <220> FEATURE:
- 79 <223> OTHER INFORMATION: Targeting Vector
- 81 <400> SEQUENCE: 3
- 82 ggatgagttg aagccagaac agcagcaccc tctcccttcc tacctggagt acagacacaa 60
- 83 cgcctccatc cgagtctacc agaccaatta ttttgccagc aaatgtgctg aaagtcccga 120
- 84 cagcagttct aaaagtttca acattaagct tggagcagag atggacagcc tcggtggcaa 180
- 85 atgtgatccc agtcagcaga 200



VERIFICATION SUMMARY

PATENT APPLICATION: US/10/005,200

DATE: 12/17/2001

TIME: 12:48:32

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\12172001\J005200.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date